RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/523,117

Source: PCT

Date Processed by STIC: 2-22-05

ENTERED

PCT

RAW SEQUENCE LISTING   DATE: 02/22/2005
PATENT APPLICATION: US/10/523,117   TIME: 08:22:32

Input Set : **A:\Seqlist.txt**
Output Set: **N:\CRF4\02222005\J523117.raw**

```
 4 <110> APPLICANT: GlaxoSmithKline Biologicals s.a.
 5       Francois-Xavier BERTHET
 6       Ralph BIEMANS
 7       Philippe DENOEL
 8       Christiane FERON
 9       Karine GORAJ
10       Jan POOLMAN
11       Vincent WEYNANTS
13 <120> TITLE OF INVENTION: Neisserial Vaccine Compositions Comprising a Combination of
Antigens
15 <130> FILE REFERENCE: B45315
C--> 17 <140> CURRENT APPLICATION NUMBER: US/10/523,117
18 <141> CURRENT FILING DATE: 2005-02-02
20 <150> PRIOR APPLICATION NUMBER: PCT/EP03/08571
21 <151> PRIOR FILING DATE: 2003-07-31
23 <150> PRIOR APPLICATION NUMBER: GB 0305028.3
24 <151> PRIOR FILING DATE: 2003-03-05
26 <150> PRIOR APPLICATION NUMBER: GB 0230170.3
27 <151> PRIOR FILING DATE: 2002-12-24
29 <150> PRIOR APPLICATION NUMBER: GB 0230168.7
30 <151> PRIOR FILING DATE: 2002-12-24
32 <150> PRIOR APPLICATION NUMBER: GB 0230164.6
33 <151> PRIOR FILING DATE: 2002-12-24
35 <150> PRIOR APPLICATION NUMBER: GB 0225531.3
36 <151> PRIOR FILING DATE: 2002-11-01
38 <150> PRIOR APPLICATION NUMBER: GB 0225524.8
39 <151> PRIOR FILING DATE: 2002-11-01
41 <150> PRIOR APPLICATION NUMBER: GB 0220199.4
42 <151> PRIOR FILING DATE: 2002-08-30
44 <150> PRIOR APPLICATION NUMBER: GB 0220197.8
45 <151> PRIOR FILING DATE: 2002-08-30
47 <150> PRIOR APPLICATION NUMBER: GB 0218037.0
48 <151> PRIOR FILING DATE: 2002-08-02
50 <150> PRIOR APPLICATION NUMBER: GB 0218036.2
51 <151> PRIOR FILING DATE: 2002-08-02
53 <150> PRIOR APPLICATION NUMBER: GB 0218035.4
54 <151> PRIOR FILING DATE: 2002-08-02
56 <150> PRIOR APPLICATION NUMBER: GB 0218051.1
57 <151> PRIOR FILING DATE: 2002-08-02
59 <160> NUMBER OF SEQ ID NOS: 24
61 <170> SOFTWARE: FastSEQ for Windows Version 4.0
63 <210> SEQ ID NO: 1
64 <211> LENGTH: 9
65 <212> TYPE: PRT
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```
 66 <213> ORGANISM: Neisseria meningitidis
 68 <220> FEATURE:
 69 <221> NAME/KEY: VARIANT
 70 <222> LOCATION: (1)...(9)
 71 <223> OTHER INFORMATION: Xaa = Any Amino Acid
 73 <400> SEQUENCE: 1
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W--> **74 Leu Xaa Gly Gly Xaa Gly Asn Asp Xaa**
```
 75  1               5
 78 <210> SEQ ID NO: 2
 79 <211> LENGTH: 9
 80 <212> TYPE: PRT
 81 <213> ORGANISM: Neisseria meningitidis
 83 <220> FEATURE:
 84 <221> NAME/KEY: VARIANT
 85 <222> LOCATION: (1)...(9)
 86 <223> OTHER INFORMATION: Xaa = Any Amino Acid
 88 <400> SEQUENCE: 2
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W--> **89 Leu Xaa Gly Gly Xaa Gly Asp Asp Xaa**
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 90  1               5
 93 <210> SEQ ID NO: 3
 94 <211> LENGTH: 10
 95 <212> TYPE: PRT
 96 <213> ORGANISM: Neisseria meningitidis
 98 <400> SEQUENCE: 3
 99 Lys Thr Lys Cys Lys Phe Leu Lys Lys Cys
100  1               5                   10
103 <210> SEQ ID NO: 4
104 <211> LENGTH: 31
105 <212> TYPE: DNA
106 <213> ORGANISM: Artificial Sequence
108 <220> FEATURE:
109 <223> OTHER INFORMATION: primer
111 <400> SEQUENCE: 4
112 ggaattccat atgatgaaca aaatataccg c                          31
114 <210> SEQ ID NO: 5
115 <211> LENGTH: 31
116 <212> TYPE: DNA
117 <213> ORGANISM: Artificial Sequence
119 <220> FEATURE:
120 <223> OTHER INFORMATION: primer
122 <400> SEQUENCE: 5
123 gtagctagct agcttaccac tgataaccga c                          31
125 <210> SEQ ID NO: 6
126 <211> LENGTH: 36
127 <212> TYPE: DNA
128 <213> ORGANISM: Artificial Sequence
130 <220> FEATURE:
131 <223> OTHER INFORMATION: primer
133 <400> SEQUENCE: 6
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134 aactgcagaa ttaatatgaa aggagaagaa cttttc                    36
136 <210> SEQ ID NO: 7
137 <211> LENGTH: 33
138 <212> TYPE: DNA
139 <213> ORGANISM: Artificial Sequence
141 <220> FEATURE:
142 <223> OTHER INFORMATION: primer
144 <400> SEQUENCE: 7
145 gacatactag tttatttgta gagctcatcc atg                        33
147 <210> SEQ ID NO: 8
148 <211> LENGTH: 30
149 <212> TYPE: DNA
150 <213> ORGANISM: Artificial Sequence
152 <220> FEATURE:
153 <223> OTHER INFORMATION: primer
155 <400> SEQUENCE: 8
156 tccccgcggg ccgtctgaat acatcccgtc                            30
158 <210> SEQ ID NO: 9
159 <211> LENGTH: 51
160 <212> TYPE: DNA
161 <213> ORGANISM: Artificial Sequence
163 <220> FEATURE:
164 <223> OTHER INFORMATION: primer
166 <400> SEQUENCE: 9
167 catatgggct tccttttgta aatttgaggg caaacacccg atacgtcttc a     51
169 <210> SEQ ID NO: 10
170 <211> LENGTH: 48
171 <212> TYPE: DNA
172 <213> ORGANISM: Artificial Sequence
174 <220> FEATURE:
175 <223> OTHER INFORMATION: primer
177 <400> SEQUENCE: 10
178 agacgtatcg ggtgtttgcc ctcaaattta caaaggaag cccatatg          48
180 <210> SEQ ID NO: 11
181 <211> LENGTH: 33
182 <212> TYPE: DNA
183 <213> ORGANISM: Artificial Sequence
185 <220> FEATURE:
186 <223> OTHER INFORMATION: primer
188 <400> SEQUENCE: 11
189 gggtattccg ggcccttcag acggcgcagc agg                        33
191 <210> SEQ ID NO: 12
192 <211> LENGTH: 45
193 <212> TYPE: DNA
194 <213> ORGANISM: Artificial Sequence
196 <220> FEATURE:
197 <223> OTHER INFORMATION: primer
199 <400> SEQUENCE: 12
200 ggcctagcta gccgtctgaa gcgattagag tttcaaaatt tattc           45
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202 <210> SEQ ID NO: 13
203 <211> LENGTH: 42
204 <212> TYPE: DNA
205 <213> ORGANISM: Artificial Sequence
207 <220> FEATURE:
208 <223> OTHER INFORMATION: primer
210 <400> SEQUENCE: 13
211 ggccaagctt cagacggcgt tcgaccgagt ttgagccttt gc                    42
213 <210> SEQ ID NO: 14
214 <211> LENGTH: 39
215 <212> TYPE: DNA
216 <213> ORGANISM: Artificial Sequence
218 <220> FEATURE:
219 <223> OTHER INFORMATION: primer
221 <400> SEQUENCE: 14
222 tcccccggga agatctggac gaaaaatctc aagaaaccg                        39
224 <210> SEQ ID NO: 15
225 <211> LENGTH: 64
226 <212> TYPE: DNA
227 <213> ORGANISM: Artificial Sequence
229 <220> FEATURE:
230 <223> OTHER INFORMATION: primer
232 <400> SEQUENCE: 15
233 ggaagatctc cgctcgagca aatttacaaa aggaagccga tatgcaacag caacatttgt 60
234 tccg                                                             64
236 <210> SEQ ID NO: 16
237 <211> LENGTH: 36
238 <212> TYPE: DNA
239 <213> ORGANISM: Artificial Sequence
241 <220> FEATURE:
242 <223> OTHER INFORMATION: primer
244 <400> SEQUENCE: 16
245 ggaagatctc cgctcgagac atcgggcaaa cacccg                           36
247 <210> SEQ ID NO: 17
248 <211> LENGTH: 38
249 <212> TYPE: DNA
250 <213> ORGANISM: Artificial Sequence
252 <220> FEATURE:
253 <223> OTHER INFORMATION: primer
255 <400> SEQUENCE: 17
256 tcccccggga gatctcacta gtattaccct gttatccc                         38
258 <210> SEQ ID NO: 18
259 <211> LENGTH: 41
260 <212> TYPE: DNA
261 <213> ORGANISM: Artificial Sequence
263 <220> FEATURE:
264 <223> OTHER INFORMATION: primer
266 <400> SEQUENCE: 18
267 ctcgagacca tgggcaaata tcatgtctac gaccccctcg c                     41
```

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269 <210> SEQ ID NO: 19
270 <211> LENGTH: 43
271 <212> TYPE: DNA
272 <213> ORGANISM: Artificial Sequence
274 <220> FEATURE:
275 <223> OTHER INFORMATION: primer
277 <400> SEQUENCE: 19
278 gtgcatagtg tcagagtttt tgtcgacgtc gtaattatag acc               43
280 <210> SEQ ID NO: 20
281 <211> LENGTH: 39
282 <212> TYPE: DNA
283 <213> ORGANISM: Artificial Sequence
285 <220> FEATURE:
286 <223> OTHER INFORMATION: primer
288 <400> SEQUENCE: 20
289 aatggaatac atatgaataa aggtttacat cgcattatc                    39
291 <210> SEQ ID NO: 21
292 <211> LENGTH: 31
293 <212> TYPE: DNA
294 <213> ORGANISM: Artificial Sequence
296 <220> FEATURE:
297 <223> OTHER INFORMATION: primer
299 <400> SEQUENCE: 21
300 ccaactagtg tttttcgcta cttggagctg t                            31
302 <210> SEQ ID NO: 22
303 <211> LENGTH: 44
304 <212> TYPE: DNA
305 <213> ORGANISM: Artificial Sequence
307 <220> FEATURE:
308 <223> OTHER INFORMATION: primer
310 <400> SEQUENCE: 22
311 aatggaatac atatgaataa aggtttacat cgcattatct ttag              44
313 <210> SEQ ID NO: 23
314 <211> LENGTH: 45
315 <212> TYPE: DNA
316 <213> ORGANISM: Artificial Sequence
318 <220> FEATURE:
319 <223> OTHER INFORMATION: primer
321 <400> SEQUENCE: 23
322 ggggccactc gaggtttttc gctacttgga gctgtttcag atagg             45
324 <210> SEQ ID NO: 24
325 <211> LENGTH: 4217
326 <212> TYPE: DNA
327 <213> ORGANISM: Neisseria meningitidis
329 <400> SEQUENCE: 24
330 atgaataaag gtttacatcg cattatcttt agtaaaaagc acagcaccat ggttgcagta 60
331 gccgaaactg ccaacagcca gggcaaaggt aaacaggcag gcagttcggt ttctgtttca 120
332 ctgaaaactt caggcgacct ttgcggcaaa ctcaaaacca cccttaaaac tttggtctgc 180
333 tctttggttt ccctgagtat ggtattgcct gcccatgccc aaattaccac cgacaaatca 240
```

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 02/22/2005
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<u>Please Note:</u>
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 2,5,9
Seq#:2; Xaa Pos. 2,5,9

VERIFICATION SUMMARY   DATE: 02/22/2005
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L:17 M:270 C: Current Application Number differs, Replaced Current Application Number
L:74 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:89 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0